

TITLE	A large scale analysis of mRNAs expressed by primary mesenchyme cells of the sea urchin embryo Development 128 (13), 2615-2627 (2001)
JOURNAL MEDLINE	21384984
COMMENT	Contact: Ettensohn CA Dept. Biol. Sci. Carnegie Mellon University 4400 Fifth Avenue, Pittsburgh, PA 15213, USA Tel.: +1 412 268 5849 Email: ettensohn@andrew.cmu.edu. Location/Qualifiers
FEATURES	source
BASE COUNT	136 a 241 c 184 g 236 t 1 others
ORIGIN	
Query Match	13.1%; Score 230.4; DB 12; Length 798;
Best Local Similarity	57.3%; Pred. No. 1e-49;
Matches 434:	Conservative 0; Mismatches 322; Indels 1; Gaps 1;
Dy	715 GTGGTGTGGGAGACGCTCTGGCCCGGACGTGGTGCTGATTCTGCGGAGAGC 774
Dd	
Oy	42 GTGGTGTACTTCACAGCACTCTCCCTTACTTTGTCTGGTGCATCTCTTGTGGTCTGTA 101
Dy	775 GTCAAGCTTCCAGAGCGAGCGAGGCGCATACGCTACTACTTACCOCAGAGTGGCACAA 834
Dd	
Oy	102 GTTACTTTCACCGGGGCTCCTATGATGGATTTTGTCTTCATCGTTCCTGACTTACTACA 161
Dd	
Oy	835 TTGGAACACTAAGTATGATGATGACGCCGCGCATCCGATTTTCTCTGCTGGTCCC 894
Dd	162 CTCAGAACCCCTCAGGATGAGTGGCTGTATGCCCTGTGCGAATCTTCTTCCCTCAGTCC 221
Oy	895 GGGTTTCGGAACCTCTACTGGCGCTCTCCAGCTACAAACAAGTTCAACAACATGTGCACAG 954
Dd	222 GCCTGGGAGGCTCATCACCCTCTCATACCTACAAACAAGTTCCACAACAACATGTGACTTT 281
Oy	955 GACCGCGCTCATCTCTTCTTATCACTGCTTGACCACTCTTCTGCTGCTGCTCAAT 1014
Dd	282 GATCTGTGCTTGTAGCAACCCCTCAACCTGTGACTATGCTTCTTCCGCTGTTTGTGATC 341
Oy	1015 TTTCTCGGTTTTGGGTTATCGGCGAGCTTCAGAACAAGACATCGAGAGAGTTGGCCTC 1074
Dd	342 TTCTCTATCGTGGGTTTTATATGCCCCAACCTCAGGGCACAAGAAGTGGCTGATCGAC 401
Oy	1075 GAAGCCCTTGACTGTGTTGATCGTGTACCCCGAGGCCATCGCCACCATGACCGGCTCC 1134
Dd	402 TCAGAGATTGTGCTTGTCTTATGCTTATGCCCCGAGGCCGCTCCCTCATGGCCGTATCA 461
Oy	1135 GTGTTTGGGCGCATCTTCTTCTCTATGCTTATTAACCTTGGGACTGCAATGACTTTT 1199
Dd	462 CCGCTGTGGCGCATCTCTTCTTCTTATGTTGCTCAACCCCTCGCTCGATCCCATGTC 521
Oy	1195 GGAGGCTTTGAGGAGTACCAACGCGCTTTTGGCGAGATATCCCGAGTGTTAGACAGA 1254
Dd	522 AACATCATGAGAGAGCGTGGTACACAGCCCTCGTCGACGAGTTCGCCGACACCTCCGCAAG 581
Oy	1255 CATCGCGAAGTATTTGGCGTACTGCTTCTGTTTCATCTATATTTTGGCTCTGCCAAC 1314
Dd	582 AAGAAGACCTTCATCATGTTGGTGGCGCTGACATCATGCTTCCGATATCACGTCG 641
Oy	1315 ACCACATCGGTTGGTGTACTCTGTACCTACTACATATGTTATGAGCCCTGGATTTGGCG 1374
Dd	642 ATCACAGAGCGCTGACCATCTAGCTGGTACGTTTGAAGGACTCTTATGAGAGCTGGGTTTCC 701

QY	1375	ATTCATTTCGCGATTTTTCGAGAGCTCCCGGCGTGTCTGTGGGTATATGGCGTGCACCGG	1434
Db	702	CTCATTTATGTTGGGCGCTTTTGGGAGCAATTGGCCCTTCCTCTATGTAAGCAAGNCGT	761
QY	1435	TTCTCTGAGATGTGAGACCATCTGTGGGCGACACC	1471
Db	762	TTC-CCATGACATTCCTACTATATATGCGGACCGCC	797
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DEFINITION	603029220/F1 NIH_MGC_114 Homo sapiens cDNA clone IMAGE:5199552 5',		
ACCESSION	B1756400		
VERSION	B1756400.1		
KEYWORDS	EST.		
SOURCE	human.		
ORGANISM	Homo sapiens		
REFERENCE	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;		
AUTHORS	Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.		
TITLE	NIH-MGC http://mhc.nci.nih.gov/		
JOURNAL	National Institutes of Health, Mammalian Gene Collection (MGC)		
COMMENT	Unpublished (1999)		
	Contact: Robert Strausberg, Ph.D.		
	Email: c9abps-f@mail.nih.gov		
	Tissue Procurement: Life Technologies, Inc.		
	cDNA Library Preparation: Life Technologies, Inc.		
	cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)		
	DNA Sequencing by: Incyte Genomics, Inc.		
	Clone distribution: MGC clone distribution information can be		
	found through the I.M.A.G.E. Consortium/LLNL at:		
	http://image.llnl.gov		
	Plate: L14M11499 row: n column: 01		
	High quality sequence stop: 816.		
FEATURES			
SOURCE	Location/Qualifiers		
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	/clone="IMAGE:5199552"		
	/clone_id="NIH_MGC_114"		
	/lab_host="DH10B"		
	/note="Organ: brain; Vector: pCMV-Sport6; Site_1: NotI;		
	Site_2: EcoRV (destroyed); RNA source anonymous pool of 6		
	male brains, age range 23-27 yo. Library is oligo-dT		
	primed and directionally cloned (EcoRV site is destroyed		
	upon cloning). Average insert size 1.5 kb, insert size		
	range 1-3 kb. Library is normalized and enriched for		
	full-length clones and was constructed by C. Gruber		
	(Invitrogen). Research Genetics tracking code 019. Note:		
	this is a NIH_MGC Library."		
BASE COUNT	148 a 251 c 221 g 216 t		
ORIGIN			
Query Match	13.1%;	Score 230.4;	DB 13; Length 836;
Best Local Similarity	61.5%;	Pred. No. 1e-49;	
Matches 369;	Conservative 0;	Mismatches 231;	Indels 0; Gaps 0;
QY	607	GGCCGTGATGATGACATGGGCGCATCAAGCCGCTGCGTGTGTTCGGGGCTCTTT	666
Db	96	GGCATCCATGACGTGGGCGCTCCCTGGCGAGCGTGGCGGCTGTGCTCTGCGCCG	155
QY	667	GTCTCTGTCTACTTCTCTTGTGGAAGAGTCAAGAGTGTGCGCAAGGTGGTGGGTG	726
Db	156	GTCATCTGCTATTTCTGTGATCTGGAAGGGGTCACAGTCCACAGGCAAGGTATATTTC	215
QY	727	ACAGTCTGGCCCGTACGTGGTGTGATTTTCGTTGGGAGAGGCGCTACGCTGTCA	786
Db	216	ACAGCGACGTTTCCGTACTATGCTTTTCATTTTGTGATCAGAGGTGTCACCTTCC	275
QY	787	GGAGGACGAGGAGGATACGCTACTACCTTACCCGAGTGGGCAAAATTCGAAACTCT	846

RESULT 10

VERSION BQ941298.1 GI:22356776

Mammalia: Eutheria: Rodentia: Sciurognathi: Muridae: Murinae: Mus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi: Muridae: Murinae: Mus

Tissue Procurement: The Cenko Laboratory

source	1.	.907
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Query Match 13.08; Score 229.2; DB 14; Length 907;

Best Local Similarity 61.3%; Pred. No. 2.2e-49;
Matches 369; Conservative 0; Mismatches 233; Indels 0; Gaps 0;

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ACCESSION      BM951745
VERSION        BM951745.1  GT:19435335

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REFERENCES
1. (bases 1 to 100),
AUTHORS
NIH-MGC <http://mac.ncl.nih.gov/>

This clone was contributed by the Brain Molecular Anatomy Project

(BMAP)
Seq primer: pYX-5.

FEATURES
source

Location/Qualifiers
1. 660
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/strain="C57BL/6"
/db_xref="taxon:10090"
/clone_image:3685362
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/tissue_type="whole brain"
/dev_stage="embryo 18.5 dpc"
/lab_host="DH10B (T1 phage resistant)"
/note="Organ: Brain; Vector: pIX-Asc; Site_1: EcoR I;
Site_2: Not I; The library was constructed according to
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
1996. Denatured mRNA was size fractionated on a 1% agarose
gel. First strand cDNA synthesis was primed with an
oligo-dT primer containing a Not I site. Double stranded
cDNA was size selected according to mRNA size fraction.
ligated with EcoR I adaptor, digested with Not I, and then
cloned directionally into pIX-Asc vector. The library tag
sequence located between the Not I site and the polyA tail
, is CAGCAGCAGC. This library was created for the
University of Iowa Mouse Brain Molecular Anatomy Project
(BMAP): 'Gene Discovery in the Developing Mouse Nervous
System', supported by National Institute of Mental Health
(NIMH), Hemlin Chin, Ph.D., program coordinator."

BASE COUNT
ORIGIN
136 a 186 c 170 g 168 t

Query Match 13.0% Score 229; DB 14; Length 660;
Best Local Similarity 62.4% Pred. No. 2.1e-49;
Matches 380; Conservative 0; Mismatches 220; Indels 9; Gaps 1;

468 GCTGCCATGACACCTGAGACAGAGTGAACACGCCGCTGTGACGCCGTCACCC 527
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52 GCTGCCCTGACACCTGAGACAGTCTTGACACCTGACACCTGACACCTACTTCC 111
||||| ||||||| ||||| ||||||| ||||| ||||| ||||| ||||| |||||
58 ACCTCAGA-----CTAATCTAATCTTCTACACCGGAGAGAGATTCTTGACAG 578
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112 CAGAGACACATCACCCTGACACCTCCATCCACGTCACCTGCTGAGAGTTTACTTGG 171
||||| ||||||| ||||| ||||||| ||||| ||||| ||||| ||||| |||||
579 TAAATATTGGAGCAGACACATCTACAGGCTGATGACATGGGCGCATCAAGCCGC 638
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
172 CCATCTCCGAGATCATCATCAAGAGGACTCCAGACCTGGGAGCATCATAGTGGCA 231
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
639 GCTGCTCTGTGTGTTGGGGCTTTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 698
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
232 GCTGCT 291
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
699 CAGAGTCTGCAAGTGTGTGGGTGACACCTGCGCCGCTAGCTGTCTGTAT 758
||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
292 CAAACAGCTGTGCAAGTGTGTGGGTGACACCTGCGCCGCTAGCTGTCTGTAT 351
||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||
759 TCTGTGGGAGAGGCTTCACCTCCAGAGCAGACGAGGAGGATACGTTACTACTTAC 818
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
352 CCTGTGTGAGGAGGAGCCCTCTCTGAGCCTGAGAGGGGTTTCTTTACTTGA 411
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
819 CCCAGAGTGGACAAATTCGAAACTCTAAGTATGATGAGCGGACATCCAGATTTT 878
||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
412 ACCCACTGAGCAAAACTTTGGAGACAGGGGTGTGGTGTATGCTGCGGCTCAGATCTT 471
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
879 CTTCGTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 938
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
472 TTTCCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 531
||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||
939 CAAACACTGCTACAGGAGCGCTCATCTCTCTATCACTGTTTACCAAGCTTCT 998
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592 CTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 651
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1059 CGAGAGAGT 1067

Db 652 GTCCGAGGT 660
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RESULT 12
BE260823 677 bp mRNA linear EST 26-OCT-2000
LOCUS 601150368F1 NIH_MGC_19 Homo sapiens cDNA clone IMAGE:3502912 5',
DEFINITION mRNA sequence.
ACCESSION BE260823.1 GI:9132437
VERSION BE260823.1
KEYWORDS EST.
SOURCE human.
ORGANISM Homo sapiens

REFERENCE
AUTHORS NIH-MGC <http://mgc.ncl.nih.gov/>.
TITLE 1 (bases 1 to 677)
JOURNAL National Institutes of Health, Mammalian Gene Collection (MGC)
COMMENT Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: ATCC

cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
Plate: L1CM175 row: h column: 17
High quality sequence stop: 581.

FEATURES
source

Location/Qualifiers
1. 677
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/clone_lib="NIH_MGC_19"
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/lab_host="DH10B (phage-resistant)"
/note="Organ: brain; Vector: pORF7; Site_1: XhoI; Site_2:
EcoRI; cDNA made by oligo-dT priming. Directionally
cloned into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Library constructed by Ling Hong
in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: This is a NIH_MGC library."

BASE COUNT
ORIGIN
129 a 187 c 185 g 176 t

Query Match 12.9% Score 227.8; DB 10; Length 677;
Best Local Similarity 63.3% Pred. No. 4.5e-49;
Matches 398; Conservative 0; Mismatches 227; Indels 4; Gaps 3;

597 CAAGTCTAAGGCTGATGACATGGGCGCATCAAGCCGCTGCTGCTGCTGCTGCTGCT 655
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1 CGAGACACCGGGATTCATGACATCGGCTGCCCGCAGAGCTCTCTCTCTCTCTCTCT 60
||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||
656 TCGGGGCTCTTCT 714
||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
61 TCGTCTGCTCAATCT 120
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715 GTGGTGTGGGATGACAGCTGTGCGCCGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 774
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121 GTGTGTGTGATACACAGCCGCTGCTTACTTCTGCTGCTGCTGCTGCTGCTGCTGCTG 180
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775 GTACGCTTTCAGAGAGCGAGGAGGAGCATATGCTACTTACCTTACCCAGAGTGGCAAA 834
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181 GTACGCTTTCAGAGAGCGAGGAGGAGCATATGCTACTTACCTTACCCAGAGTGGCA 240
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835 TTGCAAACTCTAAGTATGATGATGAGCGGAGATCCAGATTTTCTCTGCTGCTGCTGCT 894
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241 TTGAAAGAGCCAGGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 300
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OY	895	GGGTTTGGGAACCCATCAGTCGCGCTCTCCAGGTACAAACAAGTCTCAACAACAATCGCTACAGG	954
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OY	955	GACGGCGCTCATCACTCTTCTTATCAACATGCTTGACCAGCTTCCTTGCTGGTTTTGTCATTT	1014
Dd	361	GATGGCCCTGCTGACACAGCACGATCAACATGTAATCACACACTTCGTCTCTGGGTTGCCCATTC	420
OY	1015	TTCTCTGGTTTTGGGGTATCATGGCGGCACTTTCAGAACAMAGAGCATGAGAGAGTTGGCCCTC	1074
Dd	421	TTCTTCATCCCTTGGTTGTAATATGAGCCCATBAAACAAGATGCACATTTAGAGATGTGGCCACA	480
OY	1075	GAAGGCCCTGACATGCTGTATCATGTGTAAACCCCGAGGCCATGCGCACCATGACGCGGCTCC	1134
Dd	481	GAAAGAGCTGGCCATGATGTTTCATCTCTGTATCCAGAGGCCAATTTTCATACCTGTCTGGATCT	540
OY	1135	GTTGTTTGGGCGCATCATCTTCTTCCTCAATGCTTAATTACCTGGGAGCTTGACAGTACTTTT	1194
Dd	541	ACATTCTTGGGGCTGTGTTCTC--CGTCAATGCTCCCTGGGGCTGTGGGCTTGTACAGCTTCAT	598
●	1195	GGAGGTCTTGTAGGACAGTACCAACGGGCTTT	1223
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RESULT 13	BE868121	644 bp	mRNA	linear	EST 20-OCT-2000
LOCUS	BE868121				
DEFINITION	601443433971 NIH_MGC_65 Homo sapiens CDNA clone IMAGE:3847856 5', mRNA sequence.				

ACCESSION	BE668121	
VERSION	BE668121.1	GI:10316897
KEYWORDS	EST.	
SOURCE	human.	
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	Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.	
REFERENCE	1 (bases 1 to 644)	
AUTHORS	NIH-MGC http://mgc.ncl.nih.gov/	
TITLE	National Institutes of Health, Mammalian Gene Collection (MGC)	
JOURNAL	Unpublished (1999)	
COMMENT	Contact: Robert Strausberg, Ph.D.	

Email: c9apbbs-remail.nih.gov
Tissue Procurement: ATCC
cDNA Library Preparation: Life Technologies, Inc.
Genomic Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LHAM9562 row: m column: 09
High quality sequence stop: 642.
Location/Qualifiers

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Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.8 kb. Library constructed by Life
Technologies."
94 a 196 c 182 g 172 t
BASE COUNT
ORIGIN

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	Best Local Similarity	62.7%;	Pred. No. 4.9e-49;			
	Matches 371; Conservative	0;	Mismatches 219;	Indels	2;	Gaps 1;
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Db 2 TGACCCCTTGTCTGCTGGCCCTGCTGGCTGGCTGGCTACTTCTGTGTCTGGAGAGGGGCTCA 61

Qy 701 GGAGTGTCTGGCAGAGTGGTGTGGGTGACAGCTCTGGCCCCGTACGTGGTGGCTGATTC 760

Db 62 AATCCACAGGACAGATCGTGTACTTCACTGTCAATTCCTCCATGGTGGCTGGCGTC 121

Qy 761 TGTCTGGCAGAGGGCTCAACGCTTCCAGAG--CGACGGAGGGCATACGCTACTACCTTAC 818

Db 122 TGTGTGTGTCGGAGGTGGTGTCTGTGGCTGGCGCCCGTGGATGGCATCATTTACTATCCAA 181

Qy 819 CCCAGAGTGGCACAATATGGCAAAACTCTAAAGGTATGATTTAGCCGGGCACTCCAGATTTT 878

Db 182 GCTGTACTGTGTCAAAAGCTGGGGTCCCTCTACAGTGTGGATAGATGGGGAGACCCAGATTTT 241

Qy 879 CTTCGTGCTGGTCCCGGGTTCTGGAGACCCCTACTGGCGCTCTCCAGCTCAACAAGTTCAA 938

Db 242 CTTTTCCTTAGCCCATTTGGCCTGGGGGGCCCTCAAGCCCTGGGGAGCTCAACCCGCTTCAA 301

Qy 939 CAACAAGCTGTACAGGAGACGCGCTCATCTTTCTTATCAAGCTGTGGACCGACTTCTT 998

Db 302 CAACAAGCTGTACAGGAGACGCGCATCATCTTGGCTCTCAACAAGTGGAGACACTTCTT 361

Qy 999 TGTGTGTGTGTCATTTCTCGGTTTGGGTGTACATGGCGGCAAGCTCAAGACAAGACAT 1058

Db 362 TGTGTGTGTGTGTCTTCTCATCTCTGTGGGTGTATGGCTGTCAAGAGAGGGGCTGTACAT 421

Qy 1059 CGAGAGAGTTGGCTCGAAGGCCCTCGACACTGGTGTCACTGTACACCCGAGGACATTCG 1118

Db 422 CTCACAGTGGCAGAGTACAGTACAGGGCCGGGCTTGGCTTCACTCCGCTTACCCGGGGCTGTAC 481

Qy 1119 CACCATGACCGGCTCCGCTGTTTGTGGGCCATCATCTTCTCCATGCTTATTAACCTGGG 1178

Db 482 GCTGTATGCCAATGGGCCACCTGTGGGCGCCCTGTGCTTCTTCAATGCTGTGCTGTGG 541

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Db 542 TCTCGACGCGAGTTGTAGGTGTGGAGAGGCTGTATCATCCGGTCTCTCGAC 593

	RESULT	14		
B1851570	LOCUS			
	DEFINITION			
	ACCESSION			
	VERSION			
	KEYWORDS			
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	ORGANISM			
	REFERENCE			
	AUTHORS			
	TITLE			
	JOURNAL			
	COMMENT			

B1851570 959 bp mRNA linear EST 10-OCT-2001
 603378781F1 NCI_CGAP_Mam2 Mus musculus cDNA clone IMAGE:5391260 5' ,
 mRNA sequence.
 B1851570
 B1851570.1 GI:15992317
 EST.
 house mouse.
 Mus musculus
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Rodentia; Scurionathl; Muridae; Murinae; Mus.
 1 (bases 1 to 959)
 NIH-MGC http://mgc.nci.nih.gov/.
 National Institutes of Health, Mamalian Gene Collection (MGC)
 Unpublished (1999)
 Contact: Robert Strausberg, Ph.D.
 Email: csapbs@mail.nih.gov
 Tissue Procurement: Gilbert Smith, Ph.D.
 cDNA Library Preparation: Life Technologies, Inc.
 cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
 DNA Sequencing by: Incyte Genomics, Inc.
 Clone distribution: MGC clone distribution information can be
 found through the I.M.A.G.E. Consortium/LLNL at:
 http://image.llnl.gov
 Plate: LHAM1197 row: a column: 21
 High quality sequence stop: 774.

FEATURES	Location/Qualifiers
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	/strain="FVB/N-3"
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	/clone="IMAGE:5391260"
	/clone_id="NCI_CGAP_Mam2"

